-	CRF Errors Corrected by the STIC Systems Canch
	/
3 74	CRF Processing Dat: 1/24/10 Changed a file from non-ASCII to ASCII TERED CRF Processing Dat: 1/24/10 Edited by: Verified by: Verifie
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
•	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
1	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
	Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING DATE: 01/24/2002 PATENT APPLICATION: US/09/900,715 TIME: 09:17:46

Input Set : N:\jumbos\900715.txt

Output Set: N:\CRF3\01242002\I900715.raw

```
4 <110> APPLICANT: Allen, Keith D.
 6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING PROTEIN
        PHOSPHATASE 2C GENE DISRUPTIONS
10 <130> FILE REFERENCE: R-775
12 <140> CURRENT APPLICATION NUMBER: US 09/900,715
13 <141> CURRENT FILING DATE: 2001-07-06
15 <150> PRIOR APPLICATION NUMBER: US 60/216,104
16 <151> PRIOR FILING DATE: 2000-07-06
18 <150> PRIOR APPLICATION NUMBER: US 60/223,386
19 <151> PRIOR FILING DATE: 2000-08-07
21 <160> NUMBER OF SEQ ID NOS: 4
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 996
27 <212> TYPE: DNA
28 <213> ORGANISM: Mus musculus
30 <400> SEQUENCE: 1
31 gagactgcag cagaatatgt aaaatctcga ctcccagagg ctcttaagca gcaccttcag 60
32 gattatgaga aggacaaaga aaacagtgtt ctgacctacc agaccatcct cgagcagcag 120
33 atcttgtcaa ttgaccggga aatgctggaa aagttgacag tctcctatga tgaagcaggc 180
34 acaacgtgtt tgatcsctct actctcagat aaagacctca ccgtggccaa cgttggtgac 240
35 teteggggag tettgtgtga caaagatgge aatgecatee eettgtetea egateacaag 300
36 ccttaccage tgaaggaaag gaagaggata aagagagetg gtgggtteat cagetttaat 360
37 ggctcctgga gggtccaggg aatcctagcc atgtctcgat ccctgggaga ctatccactg 420
38 aaaaatctca acgtggtcat cccagaccca gacatcttga cctttgacct ggacaagctg 480
39 cagccggagt tcatgatctt ggcctcagat ggcctgtggg atgctttcag caatgaagaa 540
40 geggttegat teatcaagga gegettggat gageeecaet ttggggeeaa aageategte 600
41 ctgcagtcct tttacagagg ctgccctgac aacatcactg tcatggtggt gaagttcagg 660
42 aatagtagca aaacagaaga gcactgaacc ctgccagatc tcagctgccc caaactagag 720
43 gacteteaac atactgttet etteatgtag taaaaggtgt gggtataatt aggateatge 780
44 gtcccaacac agaaccccct tccctgatgg ccttgaatcc ctttggagta ctgagcagag 840
45 ggttggttcc cttgctgaca ccgcagaggc tgctaagttt gtgtcccccc agcccttcta 900
47 caggocatat gttttccttc ttaataatgt actttt
49 <210> SEQ ID NO: 2
50 <211> LENGTH: 228
51 <212> TYPE: PRT
52 <213> ORGANISM: Mus musculus
54 <220> FEATURE:
55 <221> NAME/KEY: VARIANT
56 <222> LOCATION: 66
57 <223> OTHER INFORMATION: Xaa = Any Amino Acid
```

59 <400> SEQUENCE: 2

RAW SEQUENCE LISTING DATE: 01/24/2002 PATENT APPLICATION: US/09/900,715 TIME: 09:17:47

Input Set : N:\jumbos\900715.txt

Output Set: N:\CRF3\01242002\I900715.raw

```
60 Glu Thr Ala Ala Glu Tyr Val Lys Ser Arg Leu Pro Glu Ala Leu Lys
   62 Gln His Leu Gln Asp Tyr Glu Lys Asp Lys Glu Asn Ser Val Leu Thr
                                       25
                   20
   64 Tyr Gln Thr Ile Leu Glu Gln Gln Ile Leu Ser Ile Asp Arg Glu Met
                                   40
   66 Leu Glu Lys Leu Thr Val Ser Tyr Asp Glu Ala Gly Thr Thr Cys Leu
   67
∰⇒ 68 Ile Xaa Leu Leu Ser Asp Lys Asp Leu Thr Val Ala Asn Val Gly Asp
   70 Ser Arg Gly Val Leu Cys Asp Lys Asp Gly Asn Ala Ile Pro Leu Ser
                                           90
   72 His Asp His Lys Pro Tyr Gln Leu Lys Glu Arg Lys Arg Ile Lys Arg
                                       105
                   100
   74 Ala Gly Gly Phe Ile Ser Phe Asn Gly Ser Trp Arg Val Gln Gly Ile
              115
                                   120
   76 Leu Ala Met Ser Arg Ser Leu Gly Asp Tyr Pro Leu Lys Asn Leu Asn
                               135
                                                   140
          130
   78 Val Val Ile Pro Asp Pro Asp Ile Leu Thr Phe Asp Leu Asp Lys Leu
   80 Gln Pro Glu Phe Met Ile Leu Ala Ser Asp Gly Leu Trp Asp Ala Phe
                                           170
   82 Ser Asn Glu Glu Ala Val Arg Phe Ile Lys Glu Arg Leu Asp Glu Pro
                   180
                                       185
   84 His Phe Gly Ala Lys Ser Ile Val Leu Gln Ser Phe Tyr Arg Gly Cys
                                   200
   86 Pro Asp Asn Ile Thr Val Met Val Val Lys Phe Arg Asn Ser Ser Lys
                               215
                                                   220
           210
   88 Thr Glu Glu His
   89 225
   92 <210> SEQ ID NO: 3
   93 <211> LENGTH: 200
   94 <212> TYPE: DNA
   95 <213> ORGANISM: Artificial Sequence
   97 <220> FEATURE:
   98 <223> OTHER INFORMATION: Targeting vector
   100 <400> SEQUENCE: 3
   101 gctggtggcc ttggcagtga cgaaggtgaa ggaggggtgc ttgtggctca gctctgttgc 60
   102 agcagaccag cttgtggtac actcacccag accggaacaa gctaacaggc tccctctgtc 120
   103 tttccaggtg ggttcatcag ctttaatggc tcctggaggg tccagggaat cctagccatg 180
   104 tctcgatccc tgggagacta
   106 <210> SEQ ID NO: 4
   107 <211> LENGTH: 200
   108 <212> TYPE: DNA
   109 <213> ORGANISM: Artificial Sequence
   111 <220> FEATURE:
   112 <223> OTHER INFORMATION: Targeting vector
   114 <400> SEQUENCE: 4
   115 tegteetgea gteettttae agaggetgee etgacaacat caetgteatg gtggtgaagt 60
```

RAW SEQUENCE LISTING

DATE: 01/24/2002

PATENT APPLICATION: US/09/900,715

TIME: 09:17:47

Input Set : N:\jumbos\900715.txt

Output Set: N:\CRF3\01242002\I900715.raw

116 tcaggaatag tagcaaaaca gaagggcact gaaccctgcc agatctcagc tgccccaaac 120 117 tagaggactc tcaacatact gttctcttca tgtagtaaaa ggtgtgggta taattaggat 180 118 catgcgtccc aacacagaac 200

VERIFICATION SUMMARY

DATE: 01/24/2002

PATENT APPLICATION: US/09/900,715

TIME: 09:17:48

Input Set : N:\jumbos\900715.txt

Output Set: N:\CRF3\01242002\I900715.raw

 $L:68 \ M:341 \ W:$ (46) "n" or "Xaa" used, for SEQ ID#:2